WHAT IS CLAIMED IS:

- 1. A method for the analysis of mixtures containing proteins, said method comprising the steps of:
- (a) reducing the disulfide bonds in the proteins of a sample, thereby providing thiol groups in cysteine-containing proteins;
 - (b) blocking free thiols with a blocking reagent in the sample;
 - (c) digesting the proteins in the sample to provide peptides;
- (d) reducing the disulfide bonds in the digested peptides, thereby providing thiol groups in cysteine-containing peptides for reaction;
- (e) reacting cysteine-containing peptides in the sample with a reagent, wherein said reagent comprises a thiol-specific reactive group which is attached to a polymer tag via a linker, wherein the linker can be differentially labeled with stable isotopes and wherein the polymer tag forms a covalent bond with the cysteine-containing peptides;
- (f) washing the polymer-bound peptides to remove non-covalently bound species;
 - (g) eluting the cysteine-containing peptides, and
- (h) subjecting the eluted peptides to quantitative mass spectrometry
 (MS) analysis.
- 2. The method according to claim 1, wherein said method further comprises the steps of:

performing steps (a) to (d) on a second sample;

reacting cysteine-containing labels in the second sample with a stable isotope-labeled form of the reagent, wherein in reacting step (e), the reagent used is a non-isotope labeled form the reagent;

mixing the peptides of the reacted sample following step (e) and the reacted second sample; and

performing steps (g) and (h) on the peptides in the mixture.

- 3. The method according to claim 1, wherein the reagent comprises a thiol-specific reactive group is selected from the group consisting of α -haloacetyl and maleimide.
- 4. The method according to claim 1, wherein the blocking reagent is methyl methane thiosulfonate.
 - 5. The method according to claim 1, wherein the reagent has the formula:

 A1 Linker A2 polymer

wherein A1 is the thiol-reactive group and A2 is an acid labile group to which the polymer is bound.

6. The method according to claim 5, wherein the acid-labile group bound to the polymer has the structure:

7. The method according to claim 5, wherein the polymer in the reagent is a polymer resin.

- 8. The method according to claim 7, wherein the polymer resin is a homopolymer or heteropolymer comprising a polymer selected from the group consisting of polystyrene and polyethylene glycol.
- 9. The method according to claim 8, wherein the linker contains a substitution of at least six hydrogen atoms with a stable isotope.
- 10. The method according to claim 9, wherein the linker contains ten stable isotopes.
- 11. The method according to claim 9, wherein the stable isotope is deuterium.
- (12) The method according to claim 1, wherein the non-isotope labeled reagent is

The method according to claim 1, wherein the isotope labeled reagent has the formula:

- 14. The method according to claim 1, wherein the eluted peptides are subjected to high-performance liquid chromatography-mass spectrometry (MS) analysis, two-dimensional liquid chromatography MS, or MS/MS analysis.
- 15. The method according to claim 1, wherein the proteins are digested using trypsin.
- A compound useful for capturing cysteine-containing peptides, which is selected from the group consisting of a thiol-specific reactive group attached to a non-biological polymer via a linker.
- 17. The compound according to claim 16, wherein the linker contains a substitution of at least six atoms with a stable isotope.
- 18. The compound according to claim 16, wherein the linker contains ten stable isotopes.
- 19. The compound according to claim 17, wherein the stable isotope is deuterium.

20. The compound according to claim 16, selected from the group consisting of

and

- 21. A reagent kit for the analysis of proteins by mass spectral analysis that comprises a compound of claim 16.
- 22. The reagent kit of claim 21 which comprises a set of substantially identical differentially labeled cysteine-tagging reagents.
- 23. The reagent kit of claim 22 further comprising one or more proteolytic enzymes for use in digestion of proteins to be analyzed.